

CRISPR-Cas9 mediated efficient PD-1 disruption on human primary T cells from cancer patients

Shu Su^{1,5}, Bian Hu^{2,5}, Jie Shao¹, Bin Shen³, Juan Du¹, Yinan Du², Jiankui Zhou², Lixia Yu¹, Lianru Zhang¹, Fangjun Chen¹, Huizi Sha¹, Lei Cheng¹, Fanyan Meng¹, Zhengyun Zou¹, Xingxu Huang^{2,4*}, Baorui Liu^{1*}

¹The Comprehensive Cancer Centre of Drum Tower Hospital, Medical School of Nanjing University & Clinical Cancer Institute of Nanjing University, Nanjing 210008, China

²MOE Key Laboratory of Model Animal for Disease Study, Model Animal Research Center of Nanjing University, National Resource Center for Mutant Mice, Nanjing 210061, China

³State Key Laboratory of Reproductive Medicine, Department of Histology and Embryology, Nanjing Medical University, Nanjing 210029, China

⁴School of Life Science and Technology, ShanghaiTech University, 100 Haike Rd., Pudong New Area, Shanghai 201210, China

⁵These authors contributed equally to this work.

*Correspondence: baoruiliu@nju.edu.cn; huangxx@shanghaitech.edu.cn

Supplementary Table 1 Oligonucleotides for generating sgRNA expression vectors

Primers No.	Primer Sequences (5' to 3')
hPD1sg1up	ACCGCGTGA CTTCCACATGAGCG
hPD1sg1down	AAACCGCTCATGTGGAAGTCACG
hPD1sg2for	ACCGCAGTTGTGTGACACGGAAG
hPD1sg2down	AAACCTTCCGTGTACACA ACTG
hPD1sg3for	ACCGACAGCGGCACCTACCTCTG
hPD1sg3down	AAACCAGAGGTAGGTGCCGCTGT
hPD1sg4for	ACCGGGCCCTGACCACGCTCATG
hPD1sg4down	AAACCATGAGCGTGGTCAGGGCC

Supplementary Table 2 Sequences of primers for dual sgRNA plasmids produce

Primers No.	Primer Sequences (5' to 3')
hPD1-Psg1	ATGCGTCTCAACCGCGTGACTTCCACATGAGCGGTTTTAGAGCTAGAAATAGCAAG
hPD1-Psg2	ATGCGTCTCGAAACCTTCCGTGTACACAACCTGCGGTGTTTCGTCTTCCACAAG
hPD1-Psg3	ATGCGTCTCAACCGACAGCGGCACCTACCTCTGGTTTTAGAGCTAGAAATAGCAAG
hPD1-Psg4	ATGCGTCTCGAAACCATGAGCGTGGTCAGGGCCCGGTGTTTCGTCTTCCACAAG

Supplementary Table 3 Sequences of primers for PCR amplification of targeting sites

Primers No.	Primer Sequences (5' to 3')	Amplicon (bp)
hPD1testF	GGACAACGCCACCTTCACCTGC	390
hPD1testR	CTACGACCTGGAGCTCCTGAT	

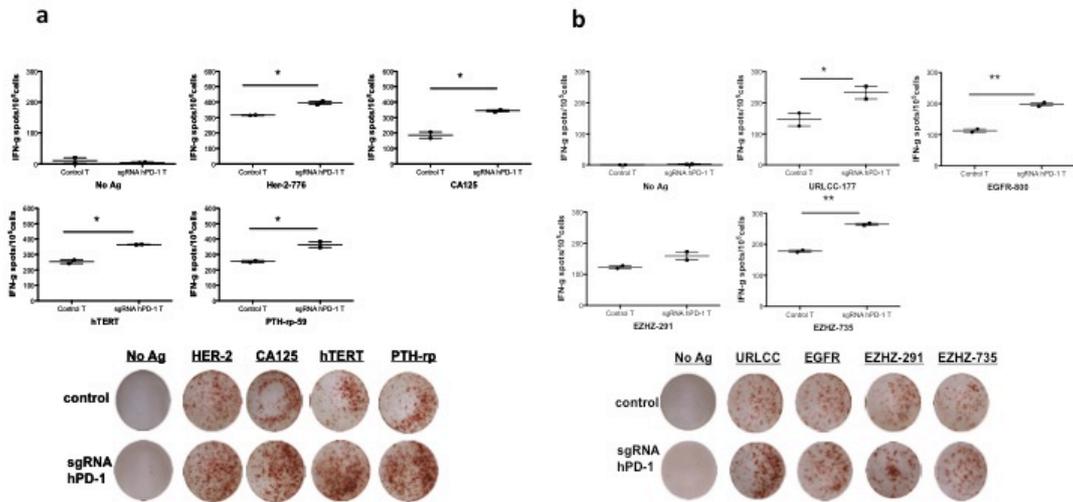
Supplementle Figure 1 Sequences of modified hPD-1 loci detected in the HeLa cellssequencing

```

hPD1
sg1  TGTCACACAACCTGCCAACGGCGTGACTTCCACATGAGCGTGGTCAGGGCCCGGGCAATGAC (WT)
      TGTCACACAACCTGCCAACGGCGGTGACTTCCACAT : AGCGTGGTCAGGGCCCGGGCAATGAC (-1, 1/21)
      TGTCACACAACCTGCCAACGGCGGTGACTTCCACAT : GCGTGGTCAGGGCCCGGGCAATGAC (-2, 2/21)
      TGTCACACAACCTGCCAACGGCGGTGACTTCCACA : GCGTGGTCAGGGCCCGGGCAATGAC (-3, 1/21)
      TGTCACACAACCTGCCAACGGCGGTGACTTCCAC : GCGTGGTCAGGGCCCGGGCAATGAC (-4, 1/21)
      TGTCACACAACCTGCCAACGGCGGTGACTTCCAC : GCGTGGTCAGGGCCCGGGCAATGAC (-8, 1/21)
      TGTCACACAACCTGCCAACGGCGG : TGAGCGTGGTCAGGGCCCGGGCAATGAC (-11, 1/21)
      TGTCACACAACCTGCCAACGGCGGTGACTTCCACATGAGCGTGGTCAGGGCCCGGGCAATGAC (+1, 6/21)
      a
sg2  CAGCCCGGCCAGGACTGCCGCTTCCGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA (WT)
      CAGCCCGGCCAGGACTGCCGCT : GTCACACAACCTGCCAACGGCGGTGACTTCCACATGA (-5, 1/19)
      CAGCCCGGCCAGGACTGCCGC : GTCACACAACCTGCCAACGGCGGTGACTTCCACATGA (-6, 1/19)
      CAGCCCGGCCAGGACTGCCGCTT : ACAACTGCCAACGGCGGTGACTTCCACATGA (-9, 1/19)
      CAGCCCGGCC : CCGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA (-13, 1/19)
      CAGCCCGGC : CGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA (-15, 1/19)
      CAGC : ACACAACCTGCCAACGGCGGTGACTTCCACATGA (-26, 1/19)
      CAGCCCGGCCAGGA : CGGCGGTGACTTCCACATGA (-30, 1/19)
      CAGCCCGGCCAGGACTGCCGCTTCCGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA (+1, 3/19)
      c
      CAGCCCGGCCAGGACTGCCGCTTCCGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA (+51, 2/19)
      bcagaaagctgcagggtttgtccataggtatacgtgtgccatggtggtttgc
sg1+2 CTGCGCCTTCCGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGAGCGTGGTCAGGGCC (WT)
      CTGCCGCTTC : AGCGTGGTCAGGGCC (-39, 1/21)
      CTGC : GCGTGGTCAGGGCC (-46, +1, 1/21)
      CTGCCGCTT : AGCGTGGTCAGGGCC (-40, +1, 2/21)
      CTGCCGCTT : gAGCGTGGTCAGGGCC (-40, +2, 1/21)
      : CGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA : AGCGTGGTCAGGGCC (-22, -9, 1/21)
      CTGCCGC : CGTGTACACAACCTGCCAACGGCGGTGACTTCCACATGA : GTGGTCAGGGCC (-3, -2, +1, 1/21)
      CTGCC : GTGTACACAACCTGCCAACGGCGGTGACTTCCACATG : tggg AGGGCC (-6, -9, +4, 1/21)
      CTGCC : GTGTACACAACCTGCCAACGGCGGTGACTTCCAC : GCGTGGTCAGGGCC (-6, -4, +1, 1/21)
      : acccggg : CGGCGGTGACTTCCACATGAGCGTGGTCAGGGCC (-25, +7, +1, 1/21)
      a
  
```

The PAM sequences are underlined and highlighted in green; the targeting sequences in red; the mutations in blue, lower case; deletions (-), insertions (+). N/N indicates positive colonies out of total sequenced.

Supplemente Figure 2 The enhanced IFN-gamma production by the Cas9-mediated disruption of PD-1 in primary T cells of cancer patients stimulated by tumor associated antigens



sgRNA hPD-1:Cas9 modified primary T cells and control T cells from cancer patients were cultured in IL-2 after electroporation for 7 d and stimulated by peptides pulsed autologous DCs for 20 h. The IFN- γ secreting cells was evaluated by Elispot. All values shown are mean \pm SD of triplicate or duplicate measurements and have been repeated 3 times with similar results. * $p < 0.05$; ** $p < 0.005$.